

## Sequence Listing

<110> Botstein,David  
Desnoyers,Luc  
Ferrara,Napoleone  
Fong,Sherman  
Gao,Wei-Qiang  
Goddard,Audrey  
Gurney,Austin L.  
Pan,James  
Roy,Margaret Ann  
Stewart,Timothy A.  
Tumas,Daniel  
Watanabe,Colin K.  
Wood,William I.

<120> Secreted and Transmembrane Polypeptides and Nucleic  
Acids Encoding the Same

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His	Val	Leu	Gly	Met	Val	Pro	Pro	Ala	Cys	Leu	Pro	Gly	Asp	Glu
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Val	Gly	Ser	Glu	Gln	Arg	Gly	Glu	Gln	Val	Thr	Asn	Gly	Arg	Glu
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Pro Glu Leu Cys	Leu Glu Glu Leu Asp Ala Ala Ile Pro Gly Ser	
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Gln Val Arg Ala	Glu Leu Ser Glu Gly Gln Arg Gln Leu Arg Glu	
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Gln Glu Phe Arg	Arg Arg Val Ala Ala Ala Gln Ser Gln Val Gln	
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Val Leu Lys Glu	Lys Lys Gln Ala Thr Glu Arg Leu Val Ser Leu	
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Ser Ala Gln Ser	Glu Lys Arg Leu Gln Glu Leu Glu Arg Asn Val	
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Gln Leu Met Arg	Gln Gln Gln Gly Gln Leu Gln Arg Arg Leu Arg	
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Glu Glu Thr Glu	Gln Lys Arg Arg Leu Glu Ala Glu Met Ser Lys	
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 Gln Ala Pro Gly Asn Glu Asp Glu Leu His Leu Ala Pro Glu Leu  
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 Leu Trp Leu Ser Pro Leu Thr Glu Gly Ala Pro Arg Thr Arg Glu  
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 Glu Thr Arg Asp Leu Val His Ala Pro Leu Pro Leu Thr Trp Lys  
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 770 775 780  
 Arg Gln Arg Glu Ala Ala Glu Pro Leu Val Gly Arg Val Leu Pro  
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 Val Gly Glu Ala Gly Leu Pro Trp Asn Phe Gly Pro Leu Ser Lys  
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 Gly Ser Gly Leu Trp Leu Cys Gln Pro Thr Pro Arg Cys Gly Asn  
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 Lys Ile Tyr Asn Pro Ser Glu Gln Cys Cys Tyr Asp Asp Ala Ile  
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 Leu Ser Leu Lys Glu Thr Arg Arg Cys Gly Ser Thr Cys Thr Phe  
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 Trp Pro Cys Phe Glu Leu Cys Cys Pro Glu Ser Phe Gly Pro Gln  
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 Gln Lys Phe Leu Val Lys Leu Arg Val Leu Gly Met Lys Ser Gln  
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 His Val Leu Tyr Pro  
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Glu	Lys	Arg	Glu	His	Ala	Thr	Arg	Asp	Gly	Pro	Gly	Arg	Val	Asn
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Glu	Leu	Gly	Arg	Pro	Ala	Arg	Asp	Glu	Gly	Gly	Ser	Gly	Arg	Asp
				65					70					75

Trp	Lys	Ser	Lys	Ser	Gly	Arg	Gly	Leu	Ala	Gly	Arg	Glu	Pro	Trp
				80					85					90

Ser	Lys	Leu	Lys	Gln	Ala	Trp	Val	Ser	Gln	Gly	Gly	Gly	Ala	Lys
				95					100					105

Ala	Gly	Asp	Leu	Gln	Val	Arg	Pro	Arg	Gly	Asp	Thr	Pro	Gln	Ala
				110					115					120

Glu	Ala	Leu	Ala	Ala	Ala	Ala	Gln	Asp	Ala	Ile	Gly	Pro	Glu	Leu
				125					130					135

Ala	Pro	Thr	Pro	Glu	Pro	Pro	Glu	Glu	Tyr	Val	Tyr	Pro	Asp	Tyr
				140					145					150

Arg	Gly	Lys	Gly	Cys	Val	Asp	Glu	Ser	Gly	Phe	Val	Tyr	Ala	Ile
				155					160					165

Gly	Glu	Lys	Phe	Ala	Pro	Gly	Pro	Ser	Ala	Cys	Pro	Cys	Leu	Cys
				170					175					180

Thr	Glu	Glu	Gly	Pro	Leu	Cys	Ala	Gln	Pro	Glu	Cys	Pro	Arg	Leu
				185					190					195

His	Pro	Arg	Cys	Ile	His	Val	Asp	Thr	Ser	Gln	Cys	Cys	Pro	Gln
				200					205					210

Cys	Lys	Glu	Arg	Lys	Asn	Tyr	Cys	Glu	Phe	Arg	Gly	Lys	Thr	Tyr
				215					220					225

Gln	Thr	Leu	Glu	Glu	Phe	Val	Val	Ser	Pro	Cys	Glu	Arg	Cys	Arg
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230	235	240
Cys Glu Ala Asn Gly Glu Val Leu Cys Thr Val Ser Ala Cys Pro		
245	250	255
Gln Thr Glu Cys Val Asp Pro Val Tyr Glu Pro Asp Gln Cys Cys		
260	265	270
Pro Ile Cys Lys Asn Gly Pro Asn Cys Phe Ala Glu Thr Ala Val		
275	280	285
Ile Pro Ala Gly Arg Glu Val Lys Thr Asp Glu Cys Thr Ile Cys		
290	295	300
His Cys Thr Tyr Glu Glu Gly Thr Trp Arg Ile Glu Arg Gln Ala		
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Met Cys Thr Arg His Glu Cys Arg Gln Met		
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 <212> DNA  
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<211> 437

<212> PRT

<213> Homo sapiens

<400> 16

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His	Val	Trp	Lys	Val	Ser	Asp	Leu	Pro	Arg	Gln	Trp	Thr	Pro	Lys
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Asn	Thr	Ser	Cys	Asp	Ser	Gly	Leu	Gly	Cys	Gln	Asp	Thr	Leu	Met
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Leu	Ile	Glu	Ser	Gly	Pro	Gln	Val	Ser	Leu	Val	Leu	Ser	Lys	Gly
				65					70					75

Cys	Thr	Glu	Ala	Lys	Asp	Gln	Glu	Pro	Arg	Val	Thr	Glu	His	Arg
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Met	Gly	Pro	Gly	Leu	Ser	Leu	Ile	Ser	Tyr	Thr	Phe	Val	Cys	Arg
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Gln	Glu	Asp	Phe	Cys	Asn	Asn	Leu	Val	Asn	Ser	Leu	Pro	Leu	Trp
				110					115					120

Ala	Pro	Gln	Pro	Pro	Ala	Asp	Pro	Gly	Ser	Leu	Arg	Cys	Pro	Val
				125					130					135

Cys	Leu	Ser	Met	Glu	Gly	Cys	Leu	Glu	Gly	Thr	Thr	Glu	Glu	Ile
				140					145					150

Cys	Pro	Lys	Gly	Thr	Thr	His	Cys	Tyr	Asp	Gly	Leu	Leu	Arg	Leu
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Arg	Gly	Gly	Gly	Ile	Phe	Ser	Asn	Leu	Arg	Val	Gln	Gly	Cys	Met
				170					175					180

Pro	Gln	Pro	Gly	Cys	Asn	Leu	Leu	Asn	Gly	Thr	Gln	Glu	Ile	Gly
				185					190					195

Pro	Val	Gly	Met	Thr	Glu	Asn	Cys	Asn	Arg	Lys	Asp	Phe	Leu	Thr
				200					205					210

Cys	His	Arg	Gly	Thr	Thr	Ile	Met	Thr	His	Gly	Asn	Leu	Ala	Gln
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Glu	Pro	Thr	Asp	Trp	Thr	Thr	Ser	Asn	Thr	Glu	Met	Cys	Glu	Val
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Thr Ser Thr Leu Val Gly Thr Lys Gly Cys Ser Thr Val Gly Ala		
260	265	270
Gln Asn Ser Gln Lys Thr Thr Ile His Ser Ala Pro Pro Gly Val		
275	280	285
Leu Val Ala Ser Tyr Thr His Phe Cys Ser Ser Asp Leu Cys Asn		
290	295	300
Ser Ala Ser Ser Ser Ser Val Leu Leu Asn Ser Leu Pro Pro Gln		
305	310	315
Ala Ala Pro Val Pro Gly Asp Arg Gln Cys Pro Thr Cys Val Gln		
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Pro Leu Gly Thr Cys Ser Ser Gly Ser Pro Arg Met Thr Cys Pro		
335	340	345
Arg Gly Ala Thr His Cys Tyr Asp Gly Tyr Ile His Leu Ser Gly		
350	355	360
Gly Gly Leu Ser Thr Lys Met Ser Ile Gln Gly Cys Val Ala Gln		
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Pro Ser Ser Phe Leu Leu Asn His Thr Arg Gln Ile Gly Ile Phe		
380	385	390
Ser Ala Arg Glu Lys Arg Asp Val Gln Pro Pro Ala Ser Gln His		
395	400	405
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 <212> DNA  
 <213> Homo sapiens

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 35 40 45  
 Tyr Phe Gly Thr Lys Thr Arg Tyr Glu Asp Val Asn Pro Val Leu  
 50 55 60  
 Leu Ser Gly Pro Glu Ala Pro Trp Arg Asp Pro Glu Leu Leu Glu  
 65 70 75  
 Gly Thr Cys Thr Pro Val Gln Leu Val Ala Leu Ile Arg His Gly  
 80 85 90  
 Thr Arg Tyr Pro Thr Val Lys Gln Ile Arg Lys Leu Arg Gln Leu  
 95 100 105  
 His Gly Leu Leu Gln Ala Arg Gly Ser Arg Asp Gly Gly Ala Ser

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Ser	Thr	Gly	Ser	Arg	Asp	Leu	Gly	Ala	Ala	Leu	Ala	Asp	Trp	Pro					
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Leu	Trp	Tyr	Ala	Asp	Trp	Met	Asp	Gly	Gln	Leu	Val	Glu	Lys	Gly					
				140					145					150					
Arg	Gln	Asp	Met	Arg	Gln	Leu	Ala	Leu	Arg	Leu	Ala	Ser	Leu	Phe					
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Pro	Ala	Leu	Phe	Ser	Arg	Glu	Asn	Tyr	Gly	Arg	Leu	Arg	Leu	Ile					
				170					175					180					
Thr	Ser	Ser	Lys	His	Arg	Cys	Met	Asp	Ser	Ser	Ala	Ala	Phe	Leu					
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Gln	Gly	Leu	Trp	Gln	His	Tyr	His	Pro	Gly	Leu	Pro	Pro	Pro	Asp					
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Val	Ala	Asp	Met	Glu	Phe	Gly	Pro	Pro	Thr	Val	Asn	Asp	Lys	Leu					
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Met	Arg	Phe	Phe	Asp	His	Cys	Glu	Lys	Phe	Leu	Thr	Glu	Val	Glu					
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Lys	Asn	Ala	Thr	Ala	Leu	Tyr	His	Val	Glu	Ala	Phe	Lys	Thr	Gly					
				245					250					255					
Pro	Glu	Met	Gln	Asn	Ile	Leu	Lys	Lys	Val	Ala	Ala	Thr	Leu	Gln					
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Val	Pro	Val	Asn	Asp	Leu	Asn	Ala	Asp	Leu	Ile	Gln	Val	Ala	Phe					
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Phe	Thr	Cys	Ser	Phe	Asp	Leu	Ala	Ile	Lys	Gly	Val	Lys	Ser	Pro					
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Trp	Cys	Asp	Val	Phe	Asp	Ile	Asp	Asp	Ala	Lys	Val	Leu	Glu	Tyr					
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Thr	Ala	Tyr	Asn	Tyr	Lys	Lys	Gln	Met	His	Arg	Lys	Phe	Arg	Ser					
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 Thr Val Ser Phe Tyr Glu Asp Leu Lys Asn His Tyr Lys Asp Ile  
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Ala	Val	Asn	Leu	Lys	Ser	Ser	Asn	Arg	Thr	Pro	Val	Val	Gln	Glu	35	40	45	
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Ser	Asp	Pro	Arg	Ile	Glu	Trp	Lys	Lys	Ile	Gln	Asp	Glu	Gln	Thr	65	70	75	
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 Arg Met Ile Arg Ser Glu Val Leu Arg Leu Val Asp Ala Ala Leu  
 65 70 75  
 Gln Asp Leu Glu Pro Gln Gln Leu Leu Leu Phe Val Gln Ser Phe  
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 Gly Ile Pro Val Ser Ser Met Ser Lys Leu Leu Gln Phe Leu Asp  
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Gln Ala Val Ala His Asp Pro Gln Thr Leu Glu Gln Asn Ile Met  
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 Asp Lys Asn Tyr Met Ala His Leu Val Glu Val Gln His Glu Arg  
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 Glu Leu Ala Arg Val Val Gln Gly Ser Pro Glu Val Pro Gly Ile  
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 350 355 360  
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Ile	His	Val	Pro	Arg	Ile	Trp	Gln	Gly	Arg	Asp	Gln	Arg	Thr	Pro
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Pro	Met	Ile	Ala	Ala	Leu	Leu	His	Gly	Arg	Thr	His	Leu	Asn	Phe	725	730	735
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His	Gln	Gly	Ala	Leu	Trp	Asp	Cys	Leu	Leu	Ser	Phe	Ile	Arg	Leu	770	775	780
Leu	Leu	Asn	Tyr	Arg	Lys	Ser	Ser	Arg	His	Leu	Ala	Ala	Phe	Ile	785	790	795
Asn	Lys	Phe	Val	Gln	Phe	Ile	His	Lys	Tyr	Ile	Thr	Tyr	Asn	Ala	800	805	810
Pro	Ala	Ala	Ile	Ser	Phe	Leu	Gln	Lys	His	Ala	Asp	Pro	Leu	His	815	820	825
Asp	Leu	Ser	Phe	Asp	Asn	Ser	Asp	Leu	Val	Met	Leu	Lys	Ser	Leu	830	835	840
Leu	Ala	Gly	Leu	Ser	Leu	Pro	Ser	Arg	Asp	Asp	Arg	Thr	Asp	Arg	845	850	855
Gly	Leu	Asp	Glu	Glu	Gly	Glu	Glu	Glu	Ser	Ser	Ala	Gly	Ser	Leu	860	865	870
Pro	Leu	Val	Ser	Val	Ser	Leu	Phe	Thr	Pro	Leu	Thr	Ala	Ala	Glu	875	880	885
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985

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&lt;400&gt; 23

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Gly	Gln	Gly	Arg	Thr	Gly	Val	Ser	Val	Val	Met	Gly	Ile	Pro	Ser	
				140					145					150	
Val	Arg	Arg	Glu	Val	His	Ser	Tyr	Leu	Thr	Asp	Thr	Leu	His	Ser	
				155					160					165	
Leu	Ile	Ser	Glu	Leu	Ser	Pro	Gln	Glu	Lys	Glu	Asp	Ser	Val	Ile	
				170					175					180	
Val	Val	Leu	Ile	Ala	Glu	Thr	Asp	Ser	Gln	Tyr	Thr	Ser	Ala	Val	
				185					190					195	
Thr	Glu	Asn	Ile	Lys	Ala	Leu	Phe	Pro	Thr	Glu	Ile	His	Ser	Gly	
				200					205					210	
Leu	Leu	Glu	Val	Ile	Ser	Pro	Ser	Pro	His	Phe	Tyr	Pro	Asp	Phe	
				215					220					225	
Ser	Arg	Leu	Arg	Glu	Ser	Phe	Gly	Asp	Pro	Lys	Glu	Arg	Val	Arg	
				230					235					240	
Trp	Arg	Thr	Lys	Gln	Asn	Leu	Asp	Tyr	Cys	Phe	Leu	Met	Met	Tyr	
				245					250					255	
Ala	Gln	Ser	Lys	Gly	Ile	Tyr	Tyr	Val	Gln	Leu	Glu	Asp	Asp	Ile	
				260					265					270	
Val	Ala	Lys	Pro	Asn	Tyr	Leu	Ser	Thr	Met	Lys	Asn	Phe	Ala	Leu	
				275					280					285	
Gln	Gln	Pro	Ser	Glu	Asp	Trp	Met	Ile	Leu	Glu	Phe	Ser	Gln	Leu	
				290					295					300	
Gly	Phe	Ile	Gly	Lys	Met	Phe	Lys	Ser	Leu	Asp	Leu	Ser	Leu	Ile	
				305					310					315	
Val	Glu	Phe	Ile	Leu	Met	Phe	Tyr	Arg	Asp	Lys	Pro	Ile	Asp	Trp	



320	325	330
Leu Leu Asp His Ile Leu Trp Val Lys Val Cys Asn Pro Glu Lys 335	340	345
Asp Ala Lys His Cys Asp Arg Gln Lys Ala Asn Leu Arg Ile Arg 350	355	360
Phe Lys Pro Ser Leu Phe Gln His Val Gly Thr His Ser Ser Leu 365	370	375
Ala Gly Lys Ile Gln Lys Leu Lys Asp Lys Asp Phe Gly Lys Gln 380	385	390
Ala Leu Arg Lys Glu His Val Asn Pro Pro Ala Glu Val Ser Thr 395	400	405
Ser Leu Lys Thr Tyr Gln His Phe Thr Leu Glu Lys Ala Tyr Leu 410	415	420
Arg Glu Asp Phe Phe Trp Ala Phe Thr Pro Ala Ala Gly Asp Phe 425	430	435
Ile Arg Phe Arg Phe Phe Gln Pro Leu Arg Leu Glu Arg Phe Phe 440	445	450
Phe Arg Ser Gly Asn Ile Glu His Pro Glu Asp Lys Leu Phe Asn 455	460	465
Thr Ser Val Glu Val Leu Pro Phe Asp Asn Pro Gln Ser Asp Lys 470	475	480
Glu Ala Leu Gln Glu Gly Arg Thr Ala Thr Leu Arg Tyr Pro Arg 485	490	495
Ser Pro Asp Gly Tyr Leu Gln Ile Gly Ser Phe Tyr Lys Gly Val 500	505	510
Ala Glu Gly Glu Val Asp Pro Ala Phe Gly Pro Leu Glu Ala Leu 515	520	525
Arg Leu Ser Ile Gln Thr Asp Ser Pro Val Trp Val Ile Leu Ser 530	535	540
Glu Ile Phe Leu Lys Lys Ala Asp 545		

&lt;210&gt; 25

&lt;211&gt; 43

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Synthetic Oligonucleotide Probe

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<211> 18

<212> DNA

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<223> Synthetic Oligonucleotide Probe

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gtccagcaag ccctcatt 18

<210> 34

<211> 20

<212> DNA

<213> Artificial Sequence

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<223> Synthetic Oligonucleotide Probe

<400> 34

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No.	Doccode	Number of pages
1	IDS	3
2	NPLI01	6
3	NPLI02	5
4	NPLI03	13

Total number of pages: 27

Remarks:

Order of re-scan issued on .....